



1351 TTGCAGTGCA GTTTCACAGA TCGTTGTTAG ATCTTTATTT TTAGCCATGC 1401 ACTGTTGTGA GGAAAAATTA CCTGTCTTGA CTGCCATGTG TTCATCATCT 1451 TAAGTATTGT AAGCTGCTAT GTATGGATTT AAACCGTAAT CATATCTTTT 1501 TCCTATCTGA GGCACTGGTG GAATAAAAA CCTGTATATT TTACTTTGTT 1551 GCAGATAGTC TTGCCGCATC TTGGCAACTT GCAGAGATGG TGGAGCTAGA 1601 AAAAAAAAA AAAAAGCCCT TTTCAGTTTG TGCACTGTGT ATGGTCCGTG 1651 TAGATTGATG CAGATTTTCT GAAATGAAAT GTTTGTTTAG ACGAGATCAT 1701 ACCGGTAAAG CAGGAATGAC AAAGCTTGCT TTTCTGGTAT GTTCTAGGTG 1751 TATTGTGACT ITTACTGTTA TATTAATTGC. CAATATAAGT AAATATAGAT 1801 TATATATGTA TAGTGTTTCA CAAAGCTTAG ACCTTTACCT TCCAGCCACC 1851 CCACAGTGCT TGATATTICA GAGTCAGTCA TTGGTTATAC ATGTGTAGTT 1901 CCAAAGCACA TAAGCTAGAA GAAGAAATAT TTCTAGGAGC ACTACCATCT 1951 GTTTTCAACA TGRAATGCCA CACACATAGA ACTCCAACAA CATCAATTTC 2001 ATTGCACAGA CTGACTGTAG TTAATTTTGT CACAGAATCT ATGGACTGAA 2051 TCTAATGCTT CCAAAAATGT TGTTTGTTTG CAAATATCAA ACATTGTTAT 2101 GCAAGAAATT ATTAATTACA AAATGAAGAT TTATACCATT GTGGTTTAAG 2151 CTGTACTGAA CTAAATCTGT GGAATGCATT GTGAACTGTA AAAGCAAAGT 2201 ATCAATAAAG CTTATAGACT TAAAAAAAA AAAAAAAAA





## SEQ ID NO:2

- 1 MEDLDQSPLV SSSDSPPRPQ PAPKYQFVRE PEDEZEEEE REEDEDEDLE
- 51 BLEVLERKYA AGLSAAPVPT APAAGAPLHD FGNDFVPFAP RGP<u>L PAAP</u>PV
- 101 APERQFSWDP SPVSSTVPAP SPLSAAAVSP SKLPEDDEPP ARPPPPPPPAS
- 151 VSPQAEPVWT FPAPAPAAPP STPAAPKRRG 8SGSVVVDLL YWRDIKKTGV
- 201 VFGASLFLLL SLTVFSIVSV TAYIALALLS VTISFRIYKG VIQAIQKSDE
- 251 GEPTRAYLES EVAISEELVO KYSNSALGEV NCTIKELRRL PLVDDLVDSL
- 301 KFAVLMWVFT YVGALFNGLT LLILALISLF SVFVIYERHQ AQIDHYLGLA
- 351 NKNVKDAMAK IQAKIPGLER KAE